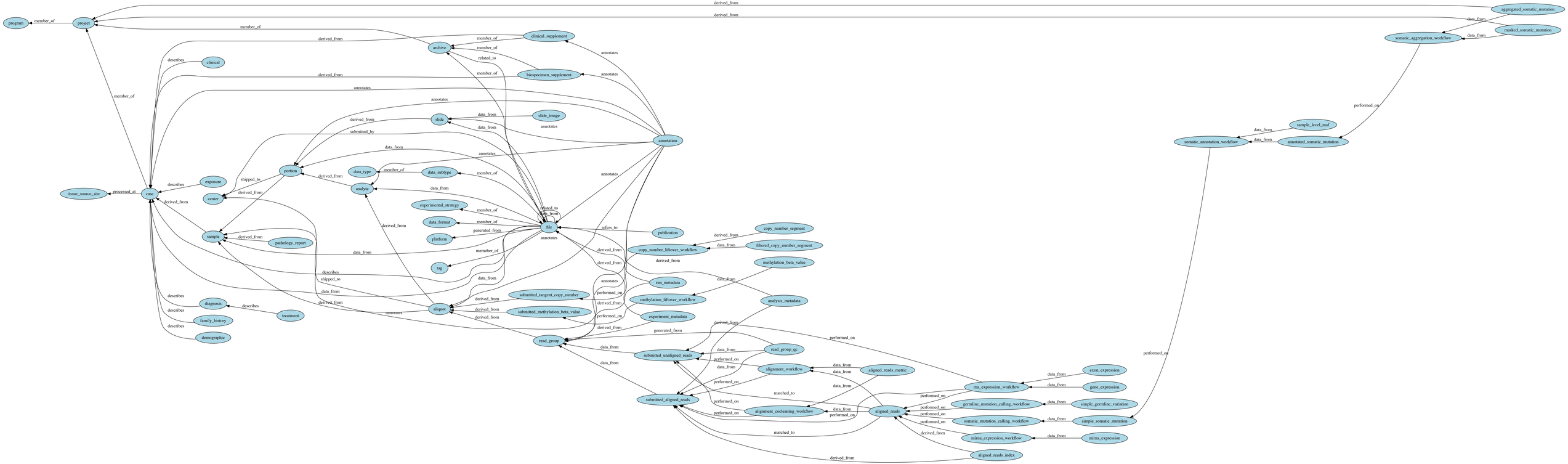
program
project
member_of
derived_from
aggregated_somatic_mutation
masked_somatic_mutation
data_from
somatic_aggregation_workflow
performed_on
sample_level_maf
somatic_annotation_workflow
annotated_somatic_mutation
archive
clinical_supplement
related_to
biospecimen_supplement
annotates
describes
clinical
slide
slide_image
submitted_by
annotation
portion
data_type
data_subtype
analyte
shipped_to
exposure
center
tissue_source_site
processed_at
case
experimental_strategy
data_format
platform
file
refers_to
publication
sample
pathology_report
tag
copy_number_segment
filtered_copy_number_segment
copy_number_liftover_workflow
methylation_beta_value
run_metadata
methylation_liftover_workflow
analysis_metadata
experiment_metadata
submitted_tangent_copy_number
submitted_methylation_beta_value
aliquot
diagnosis
treatment
family_history
demographic
read_group
read_group_qc
submitted_unaligned_reads
alignment_workflow
aligned_reads_metric
submitted_aligned_reads
alignment_cocleaning_workflow
aligned_reads
rna_expression_workflow
exon_expression
gene_expression
germline_mutation_calling_workflow
simple_germline_variation
somatic_mutation_calling_workflow
simple_somatic_mutation
mirna_expression_workflow
mirna_expression
aligned_reads_index
matched_to
generated_from
member_of
derived_from
data_from
performed_on
describes
processed_at
shipped_to
related_to
refers_to
annotates
derived_from
memeber_of